

OICE

#17

RAW SEQUENCE LISTING

DATE: 07/05/2001

PATENT APPLICATION: US/09/670,756A

TIME: 10:39:36

Input Set : A:\seqlistcorrected.txt

Output Set: N:\CRF3\07052001\I670756A.raw

3 <110> APPLICANT: Rhodes, Kenneth
 4 Betty, Maria
 5 Ling, Huai-Ping
 6 An, Wenqian
 8 <120> TITLE OF INVENTION: POTASSIUM CHANNEL INTERACTORS AND USES THEREFOR
 10 <130> FILE REFERENCE: MNI-070CP4
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/670,756A
 13 <141> CURRENT FILING DATE: 2000-09-27
 15 <150> PRIOR APPLICATION NUMBER: USSN 60/110,277
 16 <151> PRIOR FILING DATE: 1998-11-30
 18 <150> PRIOR APPLICATION NUMBER: USSN 60/110,033
 19 <151> PRIOR FILING DATE: 1998-11-25
 21 <150> PRIOR APPLICATION NUMBER: USSN 60/109,333
 22 <151> PRIOR FILING DATE: 1998-11-20
 24 <150> PRIOR APPLICATION NUMBER: USSN 09/298,731
 25 <151> PRIOR FILING DATE: 1999-04-23
 27 <150> PRIOR APPLICATION NUMBER: USSN 09/350,614
 28 <151> PRIOR FILING DATE: 1999-07-09
 30 <150> PRIOR APPLICATION NUMBER: USSN 09/350,874
 31 <151> PRIOR FILING DATE: 1999-07-09
 33 <150> PRIOR APPLICATION NUMBER: USSN 09/400,492
 34 <151> PRIOR FILING DATE: 1999-09-21
 36 <150> PRIOR APPLICATION NUMBER: USSN 09/399,913
 37 <151> PRIOR FILING DATE: 1999-09-21
 39 <150> PRIOR APPLICATION NUMBER: PCT/US99/27428
 40 <151> PRIOR FILING DATE: 1999-11-19
 43 <160> NUMBER OF SEQ ID NOS: 73
 45 <170> SOFTWARE: PatentIn Ver. 2.0
 47 <210> SEQ ID NO: 1
 48 <211> LENGTH: 1463
 49 <212> TYPE: DNA
 50 <213> ORGANISM: Homo sapiens
 52 <220> FEATURE:
 53 <221> NAME/KEY: CDS
 54 <222> LOCATION: (225)..(872)
 56 <400> SEQUENCE: 1
 58 gaatagcccc ctttcacttc tgaatccctg catgtgcggg gctqaagaag gaagccagaa 60
 61
 62
 63
 64
 65
 66
 67
 68 atg ggc acc ttc tca tct ctg caa acc aaa caa aqg cga ccc tcg aaa 284
 69 Met Gly Thr Phe Ser Ser Leu Gln Thr Lys Gln Arg Arg Pro Ser Lys
 70 5 10 15 20
 71
 72
 73
 74
 75
 76
 77
 78
 79
 80
 81
 82
 83
 84
 85
 86
 87
 88
 89
 90
 91
 92
 93
 94
 95
 96
 97
 98
 99
 100
 101
 102
 103
 104
 105
 106
 107
 108
 109
 110
 111
 112
 113
 114
 115
 116
 117
 118
 119
 120
 121
 122
 123
 124
 125
 126
 127
 128
 129
 130
 131
 132
 133
 134
 135
 136
 137
 138
 139
 140
 141
 142
 143
 144
 145
 146
 147
 148
 149
 150
 151
 152
 153
 154
 155
 156
 157
 158
 159
 160
 161
 162
 163
 164
 165
 166
 167
 168
 169
 170
 171
 172
 173
 174
 175
 176
 177
 178
 179
 180
 181
 182
 183
 184
 185
 186
 187
 188
 189
 190
 191
 192
 193
 194
 195
 196
 197
 198
 199
 200
 201
 202
 203
 204
 205
 206
 207
 208
 209
 210
 211
 212
 213
 214
 215
 216
 217
 218
 219
 220
 221
 222
 223
 224
 225
 226
 227
 228
 229
 230
 231
 232
 233
 234
 235
 236
 237
 238
 239
 240
 241
 242
 243
 244
 245
 246
 247
 248
 249
 250
 251
 252
 253
 254
 255
 256
 257
 258
 259
 260
 261
 262
 263
 264
 265
 266
 267
 268
 269
 270
 271
 272
 273
 274
 275
 276
 277
 278
 279
 280
 281
 282
 283
 284
 285
 286
 287
 288
 289
 290
 291
 292
 293
 294
 295
 296
 297
 298
 299
 300
 301
 302
 303
 304
 305
 306
 307
 308
 309
 310
 311
 312
 313
 314
 315
 316
 317
 318
 319
 320
 321
 322
 323
 324
 325
 326
 327
 328
 329
 330
 331
 332
 333
 334
 335
 336
 337
 338
 339
 340
 341
 342
 343
 344
 345
 346
 347
 348
 349
 350
 351
 352
 353
 354
 355
 356
 357
 358
 359
 360
 361
 362
 363
 364
 365
 366
 367
 368
 369
 370
 371
 372
 373
 374
 375
 376
 377
 378
 379
 380
 381
 382
 383
 384
 385
 386
 387
 388
 389
 390
 391
 392
 393
 394
 395
 396
 397
 398
 399
 400
 401
 402
 403
 404
 405
 406
 407
 408
 409
 410
 411
 412
 413
 414
 415
 416
 417
 418
 419
 420
 421
 422
 423
 424
 425
 426
 427
 428
 429
 430
 431
 432
 433
 434
 435
 436
 437
 438
 439
 440
 441
 442
 443
 444
 445
 446
 447
 448
 449
 450
 451
 452
 453
 454
 455
 456
 457
 458
 459
 460
 461
 462
 463
 464
 465
 466
 467
 468
 469
 470
 471
 472
 473
 474
 475
 476
 477
 478
 479
 480
 481
 482
 483
 484
 485
 486
 487
 488
 489
 490
 491
 492
 493
 494
 495
 496
 497
 498
 499
 500
 501
 502
 503
 504
 505
 506
 507
 508
 509
 510
 511
 512
 513
 514
 515
 516
 517
 518
 519
 520
 521
 522
 523
 524
 525
 526
 527
 528
 529
 530
 531
 532
 533
 534
 535
 536
 537
 538
 539
 540
 541
 542
 543
 544
 545
 546
 547
 548
 549
 550
 551
 552
 553
 554
 555
 556
 557
 558
 559
 560
 561
 562
 563
 564
 565
 566
 567
 568
 569
 570
 571
 572
 573
 574
 575
 576
 577
 578
 579
 580
 581
 582
 583
 584
 585
 586
 587
 588
 589
 590
 591
 592
 593
 594
 595
 596
 597
 598
 599
 600
 601
 602
 603
 604
 605
 606
 607
 608
 609
 610
 611
 612
 613
 614
 615
 616
 617
 618
 619
 620
 621
 622
 623
 624
 625
 626
 627
 628
 629
 630
 631
 632
 633
 634
 635
 636
 637
 638
 639
 640
 641
 642
 643
 644
 645
 646
 647
 648
 649
 650
 651
 652
 653
 654
 655
 656
 657
 658
 659
 660
 661
 662
 663
 664
 665
 666
 667
 668
 669
 670
 671
 672
 673
 674
 675
 676
 677
 678
 679
 680
 681
 682
 683
 684
 685
 686
 687
 688
 689
 690
 691
 692
 693
 694
 695
 696
 697
 698
 699
 700
 701
 702
 703
 704
 705
 706
 707
 708
 709
 710
 711
 712
 713
 714
 715
 716
 717
 718
 719
 720
 721
 722
 723
 724
 725
 726
 727
 728
 729
 730
 731
 732
 733
 734
 735
 736
 737
 738
 739
 740
 741
 742
 743
 744
 745
 746
 747
 748
 749
 750
 751
 752
 753
 754
 755
 756
 757
 758
 759
 760
 761
 762
 763
 764
 765
 766
 767
 768
 769
 770
 771
 772
 773
 774
 775
 776
 777
 778
 779
 780
 781
 782
 783
 784
 785
 786
 787
 788
 789
 790
 791
 792
 793
 794
 795
 796
 797
 798
 799
 800
 801
 802
 803
 804
 805
 806
 807
 808
 809
 810
 811
 812
 813
 814
 815
 816
 817
 818
 819
 820
 821
 822
 823
 824
 825
 826
 827
 828
 829
 830
 831
 832
 833
 834
 835
 836
 837
 838
 839
 840
 841
 842
 843
 844
 845
 846
 847
 848
 849
 850
 851
 852
 853
 854
 855
 856
 857
 858
 859
 860
 861
 862
 863
 864
 865
 866
 867
 868
 869
 870
 871
 872
 873
 874
 875
 876
 877
 878
 879
 880
 881
 882
 883
 884
 885
 886
 887
 888
 889
 890
 891
 892
 893
 894
 895
 896
 897
 898
 899
 900
 901
 902
 903
 904
 905
 906
 907
 908
 909
 910
 911
 912
 913
 914
 915
 916
 917
 918
 919
 920
 921
 922
 923
 924
 925
 926
 927
 928
 929
 930
 931
 932
 933
 934
 935
 936
 937
 938
 939
 940
 941
 942
 943
 944
 945
 946
 947
 948
 949
 950
 951
 952
 953
 954
 955
 956
 957
 958
 959
 960
 961
 962
 963
 964
 965
 966
 967
 968
 969
 970
 971
 972
 973
 974
 975
 976
 977
 978
 979
 980
 981
 982
 983
 984
 985
 986
 987
 988
 989
 990
 991
 992
 993
 994
 995
 996
 997
 998
 999
 1000
 1001
 1002
 1003
 1004
 1005
 1006
 1007
 1008
 1009
 1010
 1011
 1012
 1013
 1014
 1015
 1016
 1017
 1018
 1019
 1020
 1021
 1022
 1023
 1024
 1025
 1026
 1027
 1028
 1029
 1030
 1031
 1032
 1033
 1034
 1035
 1036
 1037
 1038
 1039
 1040
 1041
 1042
 1043
 1044
 1045
 1046
 1047
 1048
 1049
 1050
 1051
 1052
 1053
 1054
 1055
 1056
 1057
 1058
 1059
 1060
 1061
 1062
 1063
 1064
 1065
 1066
 1067
 1068
 1069
 1070
 1071
 1072
 1073
 1074
 1075
 1076
 1077
 1078
 1079
 1080
 1081
 1082
 1083
 1084
 1085
 1086
 1087
 1088
 1089
 1090
 1091
 1092
 1093
 1094
 1095
 1096
 1097
 1098
 1099
 1100
 1101
 1102
 1103
 1104
 1105
 1106
 1107
 1108
 1109
 1110
 1111
 1112
 1113
 1114
 1115
 1116
 1117
 1118
 1119
 1120
 1121
 1122
 1123
 1124
 1125
 1126
 1127
 1128
 1129
 1130
 1131
 1132
 1133
 1134
 1135
 1136
 1137
 1138
 1139
 1140
 1141
 1142
 1143
 1144
 1145
 1146
 1147
 1148
 1149
 1150
 1151
 1152
 1153
 1154
 1155
 1156
 1157
 1158
 1159
 1160
 1161
 1162
 1163
 1164
 1165
 1166
 1167
 1168
 1169
 1170
 1171
 1172
 1173
 1174
 1175
 1176
 1177
 1178
 1179
 1180
 1181
 1182
 1183
 1184
 1185
 1186
 1187
 1188
 1189
 1190
 1191
 1192
 1193
 1194
 1195
 1196
 1197
 1198
 1199
 1200
 1201
 1202
 1203
 1204
 1205
 1206
 1207
 1208
 1209
 1210
 1211
 1212
 1213
 1214
 1215
 1216
 1217
 1218
 1219
 1220
 1221
 1222
 1223
 1224
 1225
 1226
 1227
 1228
 1229
 1230
 1231
 1232
 1233
 1234
 1235
 1236
 1237
 1238
 1239
 1240
 1241
 1242
 1243
 1244
 1245
 1246
 1247
 1248
 1249
 1250
 1251
 1252
 1253
 1254
 1255
 1256
 1257
 1258
 1259
 1260
 1261
 1262
 1263
 1264
 1265
 1266
 1267
 1268
 1269
 1270
 1271
 1272
 1273
 1274
 1275
 1276
 1277
 1278
 1279
 1280
 1281
 1282
 1283
 1284
 1285
 1286
 1287
 1288
 1289
 1290
 1291
 1292
 1293
 1294
 1295
 1296
 1297
 1298
 1299
 1300
 1301
 1302
 1303
 1304
 1305
 1306
 1307
 1308
 1309
 1310
 1311
 1312
 1313
 1314
 1315
 1316
 1317
 1318
 1319
 1320
 1321
 1322
 1323
 1324
 1325
 1326
 1327
 1328
 1329
 1330
 1331
 1332
 1333
 1334
 1335
 1336
 1337
 1338
 1339
 1340
 1341
 1342
 1343
 1344
 1345
 1346
 1347
 1348
 1349
 1350
 1351
 1352
 1353
 1354
 1355
 1356
 1357
 1358
 1359
 1360
 1361
 1362
 1363
 1364
 1365
 1366
 1367
 1368
 1369
 1370
 1371
 1372
 1373
 1374
 1375
 1376
 1377
 1378
 1379
 1380

RAW SEQUENCE LISTING

DATE: 07/05/2001

PATENT APPLICATION: US/09/670,756A

TIME: 10:39:36

Input Set : A:\seqlistcorrected.txt

Output Set: N:\CRF3\07052001\I670756A.raw

```

74          25          30          35
76 gag gga ctg gag cag ctc gag gcc cag acc aac ttc acc aag agg gag 380
77 Glu Gly Leu Glu Gln Leu Glu Ala Gln Thr Asn Phe Thr Lys Arg Glu
78          40          45          50
80 ctg cag gtc ctt tat cga ggc ttc aaa aat gag tgc ccc agt ggt gtg 428
81 Leu Gln Val Leu Tyr Arg Gly Phe Lys Asn Glu Cys Pro Ser Gly Val
82          55          60          65
84 gtc aac gaa gac aca ttc aag cag atc tat gct cag ttt ttc cct cat 476
85 Val Asn Glu Asp Thr Phe Lys Gln Ile Tyr Ala Gln Phe Phe Pro His
86          70          75          80
88 gga gat gcc agc acg tat gcc cat tac ctc ttc aat gcc ttc gac acc 524
89 Gly Asp Ala Ser Thr Tyr Ala His Tyr Leu Phe Asn Ala Phe Asp Thr
90 85          90          95          100
92 act cag aca ggc tcc gtg aag ttc gag gac ttt gta acc gct ctg tcg 572
93 Thr Gln Thr Gly Ser Val Lys Phe Glu Asp Phe Val Thr Ala Leu Ser
94          105          110          115
96 att tta ttg aga gga act gtc cac gag aaa cta agg tgg aca ttt aat 620
97 Ile Leu Leu Arg Gly Thr Val His Glu Lys Leu Arg Trp Thr Phe Asn
98          120          125          130
100 ttg tat gac atc aac aag gac gga tac ata aac aaa gag gag atg atg 668
101 Leu Tyr Asp Ile Asn Lys Asp Gly Tyr Ile Asn Lys Glu Glu Met Met
102          135          140          145
104 gac att gtc aaa gcc atc tat gac atg atg ggg aaa tac aca tat cct 716
105 Asp Ile Val Lys Ala Ile Tyr Asp Met Met Gly Lys Tyr Thr Tyr Pro
106          150          155          160
108 gtg ctc aaa gag gac act cca agg cag cat gtg gac gtc ttc ttc cag 764
109 Val Leu Lys Glu Asp Thr Pro Arg Gln His Val Asp Val Phe Phe Gln
110 165          170          175          180
112 aaa atg gac aaa aat aaa gat ggc atc gta act tta gat gaa ttt ctt 812
113 Lys Met Asp Lys Asn Lys Asp Gly Ile Val Thr Leu Asp Glu Phe Leu
114          185          190          195
116 gaa tca tgt cag gag gac gac aac atc atg agg tct ctc cag ctg ttt 860
117 Glu Ser Cys Gln Glu Asp Asp Asn Ile Met Arg Ser Leu Gln Leu Phe
118          200          205          210
120 caa aat gtc atg taactggtga cactcagcca ttcagctctc agagacattg 912
121 Gln Asn Val Met
122          215
124 tactaaacaa ccaccttaac accctgatct gcccttgctt tgattttaca caccaactct 972
126 tgggacagaa acacctttta cactttggaa gaattctctg ctgaagactt tcttatggaa 1032
128 ccagcatca tgtggtcag tctctgattg ccaactcttc ctctttcttc tcttgagag 1092
130 agacaagatg aaatttgagt ttgttttggg agcatgctca tctctcaca ctgctgccct 1152
132 atggaaggtc cctctgctta agcttaacaa gtagtgcaca aaatatgctg cttacgtgcc 1212
134 atggaaggtc cctctgctta agcttaacaa gtagtgcaca aaatatgctg cttacgtgcc 1212
136 atggaaggtc cctctgctta agcttaacaa gtagtgcaca aaatatgctg cttacgtgcc 1212
138 atggaaggtc cctctgctta agcttaacaa gtagtgcaca aaatatgctg cttacgtgcc 1212
140 atggaaggtc cctctgctta agcttaacaa gtagtgcaca aaatatgctg cttacgtgcc 1212
142 aaactgcccc q 1463
145 <210> SEQ ID NO: 2
146 <211> LENGTH: 216

```

RAW SEQUENCE LISTING

DATE: 07/05/2001

PATENT APPLICATION: US/09/670,756A

TIME: 10:39:36

Input Set : A:\seqlistcorrected.txt

Output Set: N:\CRF3\07052001\I670756A.raw

```

147 <212> TYPE: PRT
148 <213> ORGANISM: Homo sapiens
150 <400> SEQUENCE: 2
151 Met Gly Ala Val Met Gly Thr Phe Ser Ser Leu Gln Thr Lys Gln Arg
152 1 5 10 15
154 Arg Pro Ser Lys Asp Lys Ile Glu Asp Glu Leu Glu Met Thr Met Val
155 20 25 30
157 Cys His Arg Pro Glu Gly Leu Glu Gln Leu Glu Ala Gln Thr Asn Phe
158 35 40 45
160 Thr Lys Arg Glu Leu Gln Val Leu Tyr Arg Gly Phe Lys Asn Glu Cys
161 50 55 60
163 Pro Ser Gly Val Val Asn Glu Asp Thr Phe Lys Gln Ile Tyr Ala Gln
164 65 70 75 80
166 Phe Phe Pro His Gly Asp Ala Ser Thr Tyr Ala His Tyr Leu Phe Asn
167 85 90 95
169 Ala Phe Asp Thr Thr Gln Thr Gly Ser Val Lys Phe Glu Asp Phe Val
170 100 105 110
172 Thr Ala Leu Ser Ile Leu Leu Arg Gly Thr Val His Glu Lys Leu Arg
173 115 120 125
175 Trp Thr Phe Asn Leu Tyr Asp Ile Asn Lys Asp Gly Tyr Ile Asn Lys
176 130 135 140
178 Glu Glu Met Met Asp Ile Val Lys Ala Ile Tyr Asp Met Met Gly Lys
179 145 150 155 160
181 Tyr Thr Tyr Pro Val Leu Lys Glu Asp Thr Pro Arg Gln His Val Asp
182 165 170 175
184 Val Phe Phe Gln Lys Met Asp Lys Asn Lys Asp Gly Ile Val Thr Leu
185 180 185 190
187 Asp Glu Phe Leu Glu Ser Cys Gln Glu Asp Asp Asn Ile Met Arg Ser
188 195 200 205
190 Leu Gln Leu Phe Gln Asn Val Met
191 210 215
194 <210> SEQ ID NO: 3
195 <211> LENGTH: 1856
196 <212> TYPE: DNA
197 <213> ORGANISM: Rattus sp.
199 <220> FEATURE:
200 <221> NAME/KEY: CDS
201 <222> LOCATION: (300)..(1034)
203 <400> SEQUENCE: 3
204 ggcacacaaac ccttggattc ttgggagaat atgccgtgag gtgttgccaa ttattagttc 60
206 tcttggctag cagatgttta gggactggtt aagccttgg agaaattacc ttaggaaaac 120
208 ggggaaataa aagcaaatat taccatgaat tcaagatta cctagcaatt gcaaggtatg 180
210 ggggaaataa aagcaaatat taccatgaat tcaagatta cctagcaatt gcaaggtatg 240
212 ggggaaataa aagcaaatat taccatgaat tcaagatta cctagcaatt gcaaggtatg 300
214 ggggaaataa aagcaaatat taccatgaat tcaagatta cctagcaatt gcaaggtatg 360
215 Met Pro Ala Arg Val Leu Leu Ala Tyr Pro Gly Thr Glu Met Leu Thr
216 1 5 10 15
218 cag ggc gag tct gaa ggg ctc cag acc ttg ggg ata gta gtg gtc ctg 395
219 Gln Gly Glu Ser Glu Gly Leu Gln Thr Leu Gly Ile Val Val Val Leu

```

RAW SEQUENCE LISTING

DATE: 07/05/2001

PATENT APPLICATION: US/09/670,756A

TIME: 10:39:36

Input Set : A:\seqlistcorrected.txt

Output Set: N:\CRF3\07052001\I670756A.raw

```

220          20          25          30
222 tgt tcc tct ctg aaa cta ctg cac tac ctc ggg ctg att gac ttg tcg 443
223 Cys Ser Ser Leu Lys Leu Leu His Tyr Leu Gly Leu Ile Asp Leu Ser
224          35          40          45
226 gat gac aag atc gag gat gat ctg gag atg acc atg gtt tgc cat cgg 491
227 Asp Asp Lys Ile Glu Asp Asp Leu Glu Met Thr Met Val Cys His Arg
228          50          55          60
230 cct gag gga ctg gag cag ctt gag gca cag acg aac ttc acc aag aga 539
231 Pro Glu Gly Leu Glu Gln Leu Glu Ala Gln Thr Asn Phe Thr Lys Arg
232 65          70          75          80
234 gaa ctg caa gtc ctt tac cgg gga ttc aaa aac gag tgc ccc agt ggt 587
235 Glu Leu Gln Val Leu Tyr Arg Gly Phe Lys Asn Glu Cys Pro Ser Gly
236          85          90          95
238 gtg gtt aac gaa gag aca ttc aag cag atc tac gct cag ttt ttc cct 635
239 Val Val Asn Glu Glu Thr Phe Lys Gln Ile Tyr Ala Gln Phe Phe Pro
240          100          105          110
242 cat gga gat gcc agc aca tac gca cat tac ctc ttc aat gcc ttc gac 683
243 His Gly Asp Ala Ser Thr Tyr Ala His Tyr Leu Phe Asn Ala Phe Asp
244          115          120          125
246 acc acc cag aca ggc tct qta aag ttc gag gac ttt gtg act gct ctg 731
247 Thr Thr Gln Thr Gly Ser Val Lys Phe Glu Asp Phe Val Thr Ala Leu
248          130          135          140
250 tcg att tta ctg aga gga acg gtc cat gaa aaa ctg agg tgg acg ttt 779
251 Ser Ile Leu Leu Arg Gly Thr Val His Glu Lys Leu Arg Trp Thr Phe
252 145          150          155          160
254 aat ttg tac gac atc aat aaa gac ggc tac ata aac aaa gag gag atg 827
255 Asn Leu Tyr Asp Ile Asn Lys Asp Gly Tyr Ile Asn Lys Glu Glu Met
256          165          170          175
258 atg gac ata gtg aaa gcc atc tat gac atg atg ggg aaa tac acc tat 875
259 Met Asp Ile Val Lys Ala Ile Tyr Asp Met Met Gly Lys Tyr Thr Tyr
260          180          185          190
262 cct gtg ctc aaa gag gac act ccc agg cag cac gtg gac gtc ttc ttc 923
263 Pro Val Leu Lys Glu Asp Thr Pro Arg Gln His Val Asp Val Phe Phe
264          195          200          205
266 cag aaa atg gat aaa aat aaa gat ggc att qta acg tta gac qaa ttt 971
267 Gln Lys Met Asp Lys Asn Lys Asp Gly Ile Val Thr Leu Asp Glu Phe
268          210          215          220
270 ctc gag tcc tgt cag gag gat gac aac atc atg agg tct cta cag ctg 1019
271 Leu Glu Ser Cys Gln Glu Asp Asp Asn Ile Met Arg Ser Leu Gln Leu
272 225          230          235          240
274 ttc caa aat gtc atg taactgagga cactggccat cctgctctca gagacactga 1074
275 Phe Gln Asn Val Met

```

```

182 cgaatggctc aattctctgat tggcaactct tcttccctcc tctcttggag aaggaagagc 1254
184 tgaatccga agtttgtttt qgaagcatgc ccattctctc atgctgtgc tgcctgtgg 1314
186 aaggcccttc tcttgagct taaacagtat tgcacagttt tctgcgtata cagatcccca 1374
188 actcaactgcc tetaaqtcaq qcagaccctg atcaatctga accaaatgtg caccatccctc 1434

```


VERIFICATION SUMMARY

DATE: 07/05/2001

PATENT APPLICATION: US/09/670,756A

TIME: 10:39:37

Input Set : A:\seqlistcorrected.txt

Output Set: N:\CRF3\07052001\I670756A.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application Number
L:877 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:878 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:936 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:3017 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37
L:3353 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43
L:3356 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43